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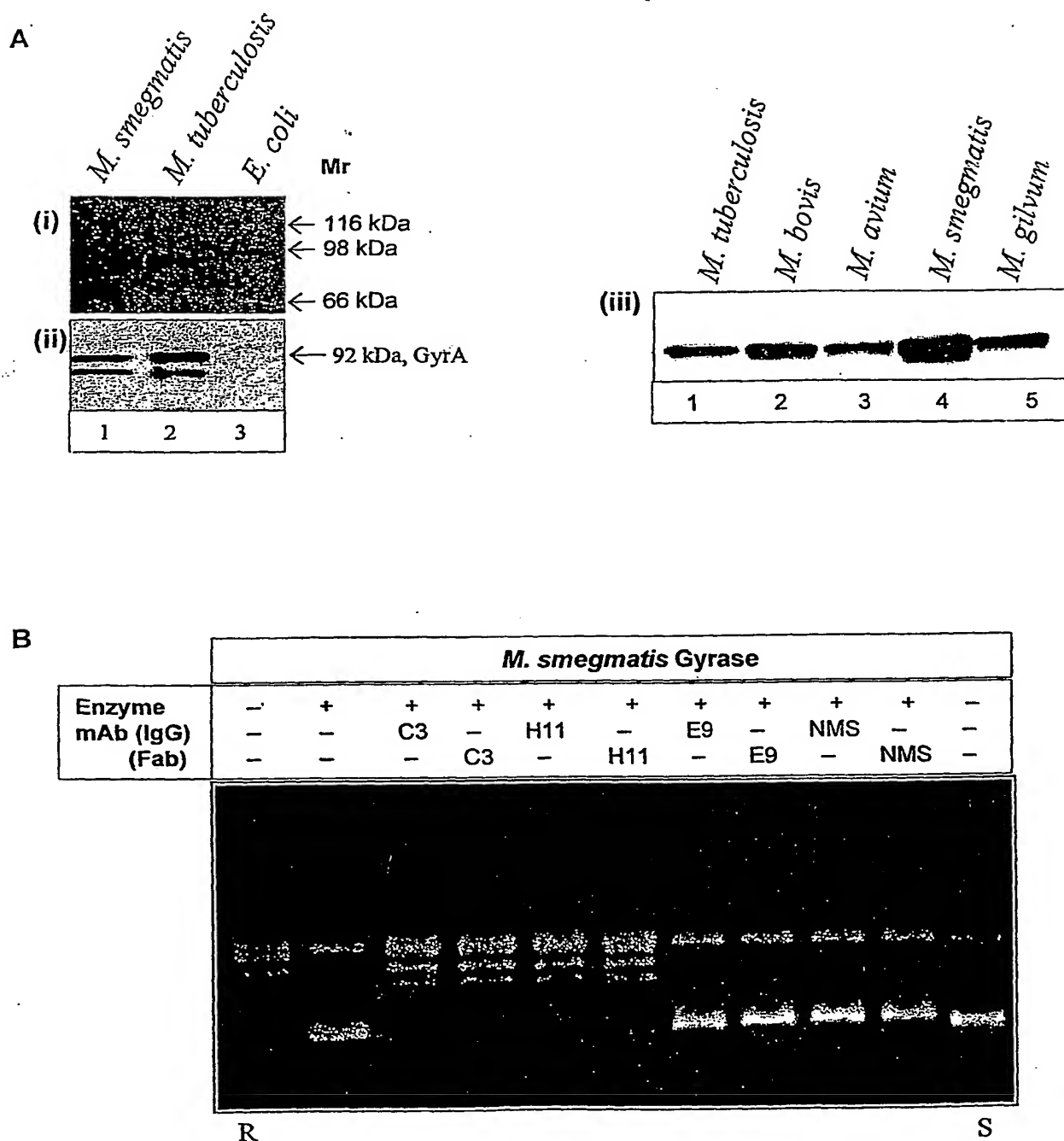


Fig. 1

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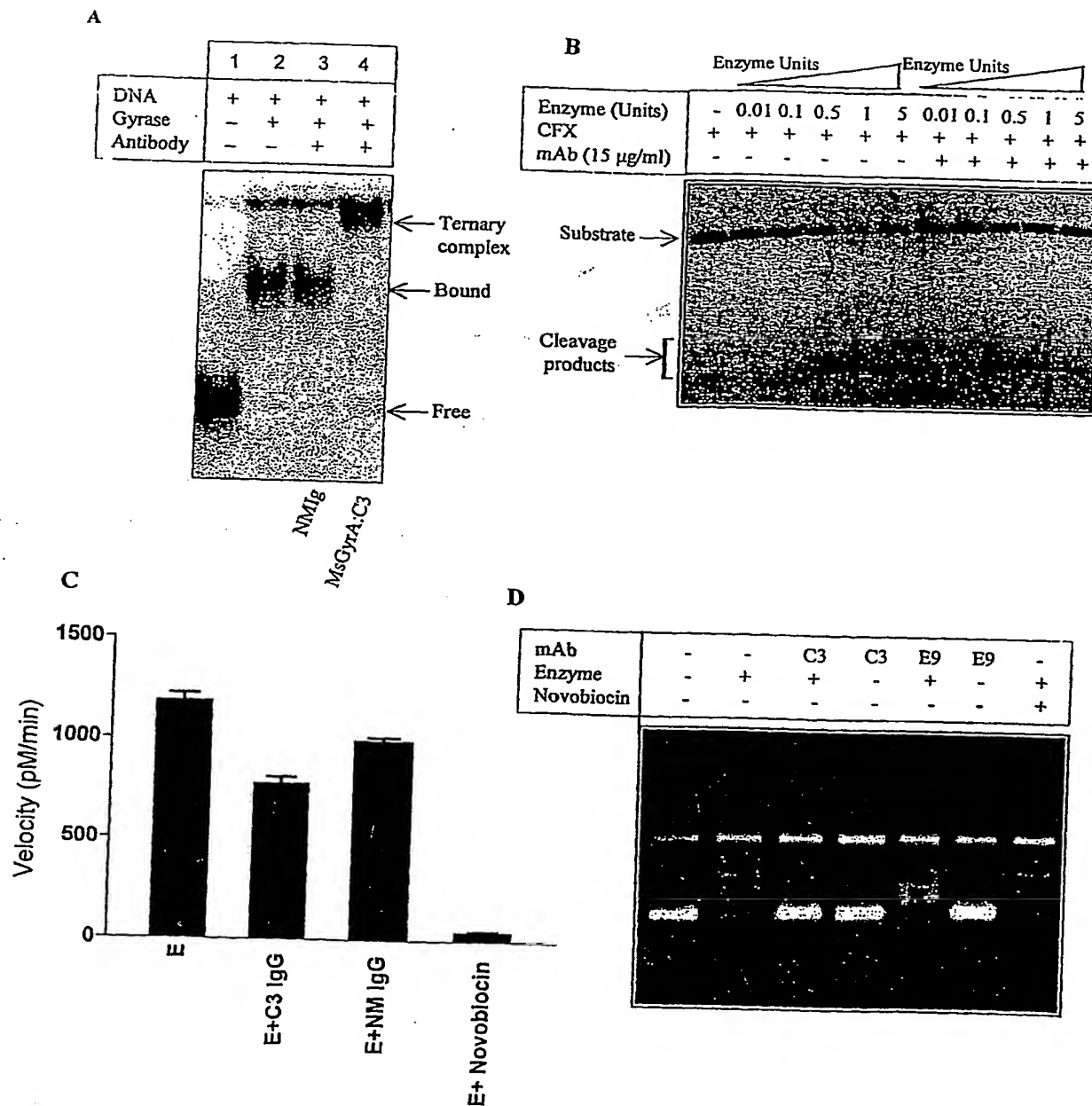


Fig. 2

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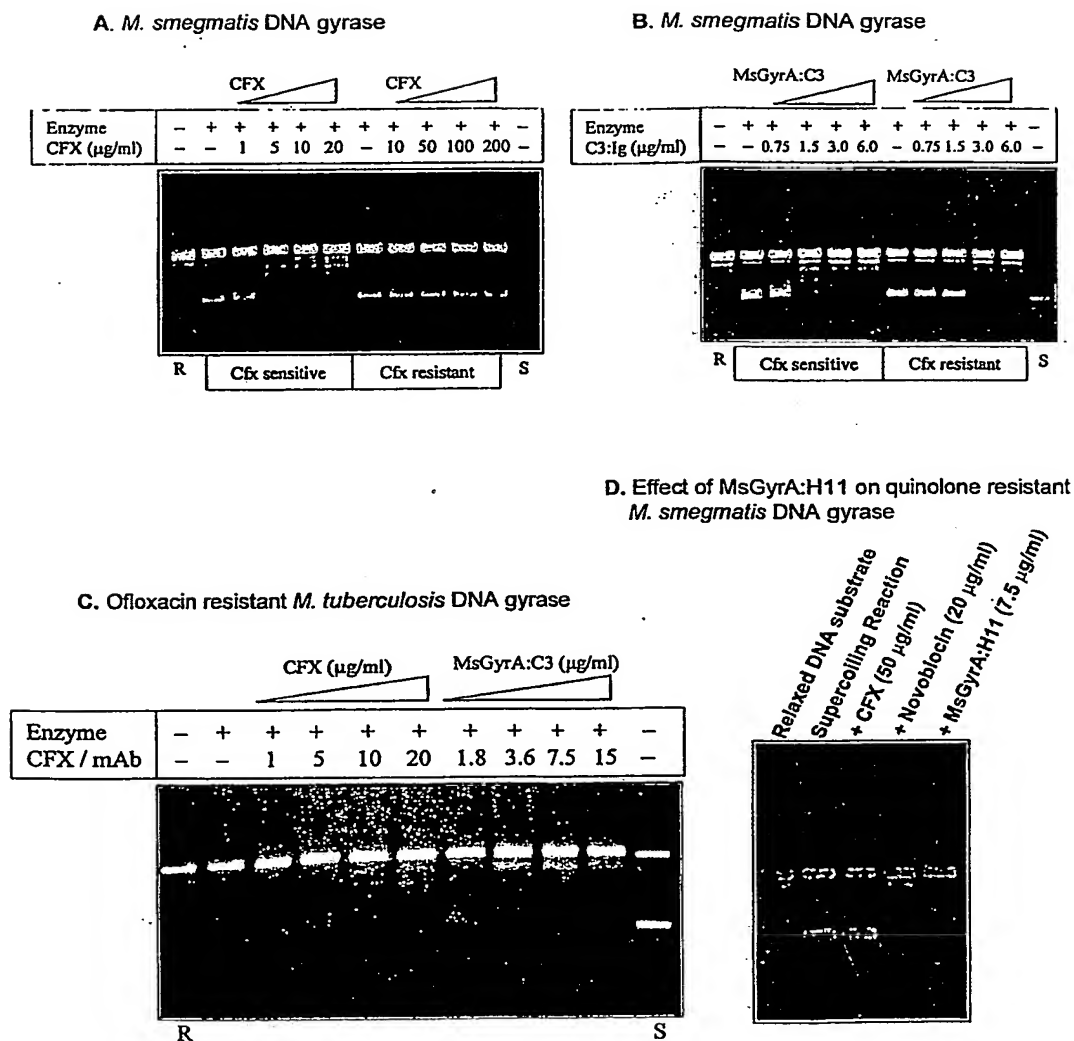


Fig. 3

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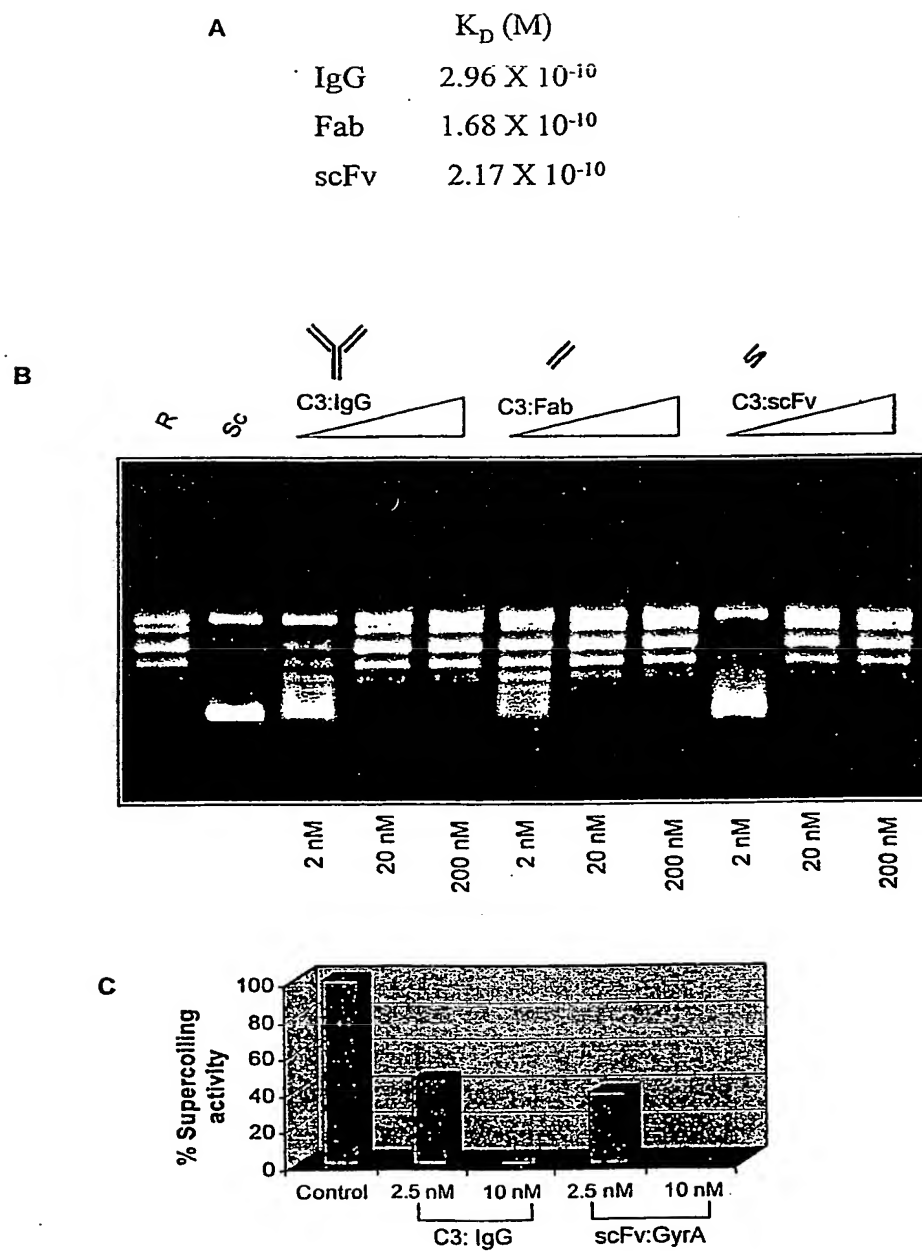


Fig. 4

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1 - GCCCAGGTGAAACTGCAGCAGTCTGGGGCTGAATTGGTGAGGCCTGGGGCTTCAGTGAAG -60  
- A Q V K L Q Q S G A E L V R P G A S V K

61 - TTGTCCTGCAAGGCTTCTGGCTACAGCTTCACCGTCTACTATATTTACTGGGTGAAACAG -120  
- L S C K A S G Y S F T V Y Y I Y W V K Q

121- AGGCCTGGACAAGCCCTTGAGTGGATTGGAGAGATTAATCCTAGCAATGGTGGTACTAAC -180  
- R P G Q A L E W I G E I N P S N G G T N

181- TTCAATGAAAAGTTCAAGACCAAGGCCACACTGACTGTAGACAAATCCACCAGCACAGTC -240  
- F N E K F K T K A T L T V D K S T S T V

241- TACATGCAACTCAGCAGCCTGACATCTGAAGACTCTGCGGTCTATTACTGTACAAGATGG -300  
- Y M Q L S S L T S E D S A V Y Y C T R W

301- GGGTTACGACGAGGGTTTGCTTACTGGGGCCAAGGGACCACGGTCACCGTCTCCTCAAGT -360  
- G L R R G F A Y W G Q G T T V T V S S S

361- GGAGGCGGTTTCAGGCGGAGGTGGCTCTGGCGGTGGCGGATCGGACATCGAGCTCACTCAG -420  
- G G G S G G G G S G G G G S D I E L T Q

421- TCTCCAAAATCCATGTCCATGTCAGTAGGAGAGAGGGTCACCTTGAGTTGCAAGGCCAGT -480  
- S P K S M S M S V G E R V T L S C K A S

481- GAGAATGTGGGTACTCATGTATCCTGGTATCAACAGAGACCAGAGGAGTCTCCTAAACTG -540  
- E N V G T H V S W Y Q Q R P E E S P K L

541- CTGATATACGGGGCATCCAACCGGTACACTGGGGTCCCCGATCGCTTCACAGGCAGTGGC -600  
- L I Y G A S N R Y T G V P D R F T G S G

601- TCTGCAACAGATTTCACTCTGACCATCAGCAATGTGCAGGCTGAAGACCTTGCAGATTAT -660  
- S A T D F T L T I S N V Q A E D L A D Y

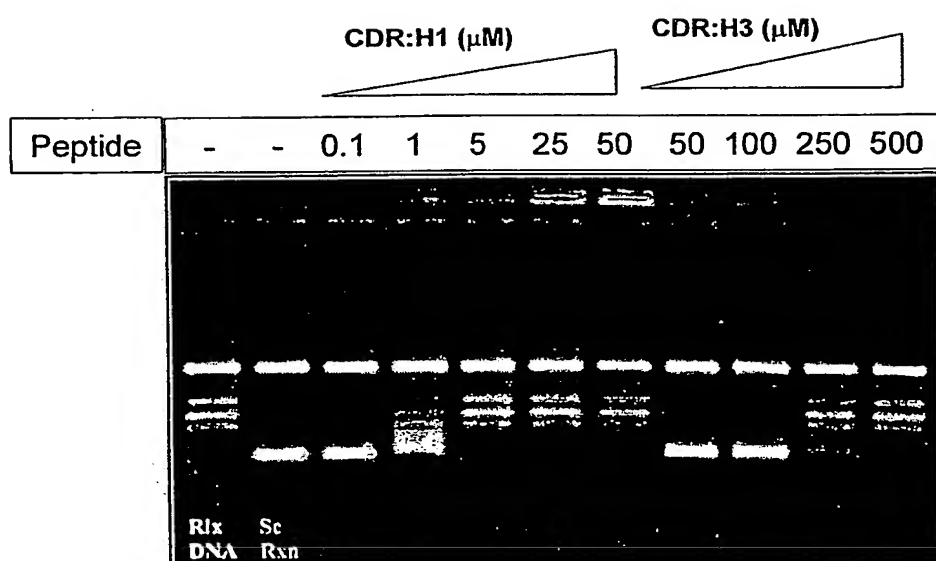
661- CACTGTGGACAGACTTACAGCTATCCATTACATTTCGGCTTGGGGACAAAGTTGGAAATA -720  
- H C G Q T Y S Y P F T F G L G T K L E I

721- AAACGGGCGGCCGAGGTGCGCCGGTGCCGTATCCGGATCCGCTGGAACCGCGTGCCGCA -780  
- K R A A A G A P V P Y P D P L E P R A A

781- TAG -783  
- \*

Fig. 5

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CDR:H1 - KASGYSFTVYYIYWVK

CDR:H3 - TRWGLRRGFAYWGQ

Fig. 6

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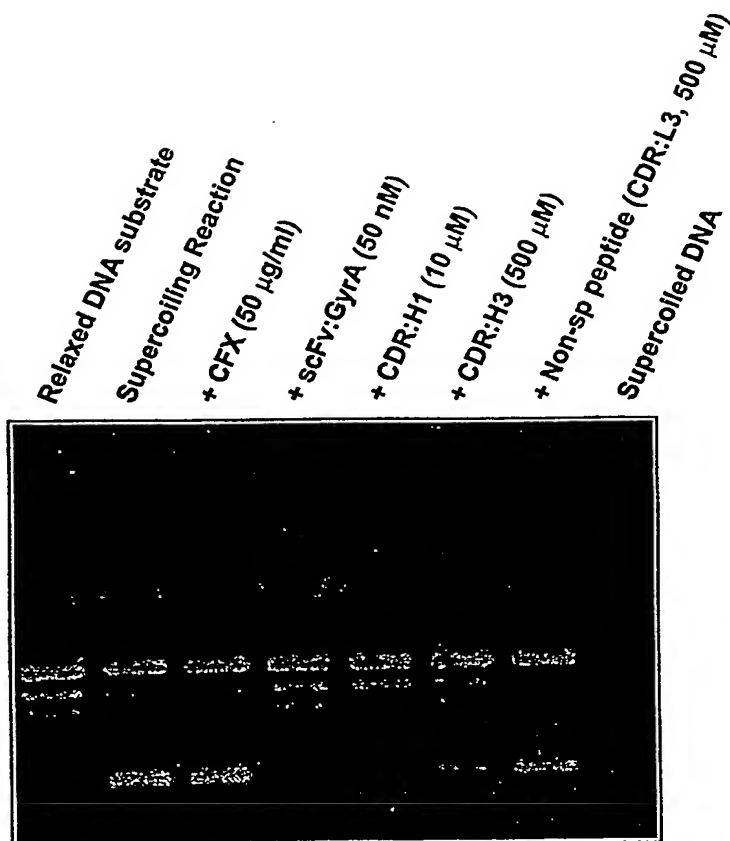


Fig. 7